OIPE

RAW SEQUENCE LISTING   DATE: 03/13/2002
PATENT APPLICATION: US/09/835,976A   TIME: 11:47:11

Input Set : A:\EP.txt
Output Set: N:\CRF3\03132002\I835976A.raw

Does Not Comply
Corrected Diskette Needed

```
   3 <110> APPLICANT: Mount, David B.
   4         Delpire, Eric
   5         Gamba, Gerardo
   6         Alfred L. George, Jr.
   8 <120> TITLE OF INVENTION: PURIFIED AND ISOLATED POTASSIUM-CHLORIDE COTRANSPORTER
NUCLEIC ACIDS AND
   9         POLYPEPTIDES AND
  10         THERAPEUTIC AND SCREENING METHODS USING SAME
  12 <130> FILE REFERENCE: Attorney Docket No. 1242-26-2
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/835,976A
C--> 15 <141> CURRENT FILING DATE: 2001-04-16
  17 <160> NUMBER OF SEQ ID NOS: 131
  19 <170> SOFTWARE: PatentIn Ver. 2.1
```

ERRORED SEQUENCES

```
4075 <210> SEQ ID NO: 14
4076 <211> LENGTH: 1083
4077 <212> TYPE: PRT
4078 <213> ORGANISM: mouse
4080 <220> FEATURE:
4081 <221> NAME/KEY: misc_feature
4082 <222> LOCATION: (71)
4083 <223> OTHER INFORMATION: Xaa=Leu or Ile
4085 <220> FEATURE:
4086 <221> NAME/KEY: misc_feature
4087 <222> LOCATION: (467)
4088 <223> OTHER INFORMATION: Xaa=Leu or Ile
4090 <220> FEATURE:
4091 <221> NAME/KEY: misc_feature
4092 <222> LOCATION: (639)
4093 <223> OTHER INFORMATION: Xaa=Leu or Ile
4095 <400> SEQUENCE: 14
4096 Met Pro Thr Asn Phe Thr Val Val Pro Val Glu Ala Arg Ala Asp Gly
4097  1               5                  10                  15
4099 Ala Gly Asp Glu Ala Ala Glu Arg Thr Glu Glu Pro Glu Ser Pro Glu
4100             20                  25                  30
4102 Ser Val Asp Gln Thr Ser Pro Thr Pro Gly Asp Gly Asn Pro Arg Glu
4103         35                  40                  45
4105 Asn Ser Pro Phe Ile Asn Asn Val Glu Val Glu Arg Glu Ser Tyr Phe
4106         50                  55                  60
4108 Glu Gly Lys Asn Met Ala Xaa Phe Glu Glu Glu Met Asp Ser Asn Pro
4109 65                  70                  75                  80
```

[handwritten notes in right margin:]

pp 1-4.

what about Xaa at location 155? (p.2)

what about Xaa at location 586? (p.3)

what about Xaa at location 1023? (p.4)

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```
        4111 Met Val Ser Ser Leu Leu Asn Lys Leu Ala Asn Tyr Thr Asn Leu Ser
        4112             85                  90                  95
        4114 Gln Gly Val Val Glu His Glu Glu Asp Glu Asp Ser Arg Arg Arg Glu
        4115             100             105             110
        4117 Val Lys Ala Pro Arg Met Gly Thr Phe Ile Gly Val Tyr Leu Pro Cys
        4118             115             120             125
        4120 Leu Gln Asn Ile Leu Gly Val Ile Leu Phe Leu Arg Leu Thr Trp Ile
        4121         130             135             140
W-->    4123 Val Gly Ala Ala Gly Val Met Glu Ser Phe Xaa Ile Val Ala Met Cys
        4124 145             150             155             160
        4126 Cys Thr Cys Thr Met Leu Thr Ala Ile Ser Met Ser Ala Ile Ala Thr
        4127             165             170             175
        4129 Asn Gly Val Val Pro Ala Gly Gly Ser Tyr Tyr Met Ile Ser Arg Ser
        4130             180             185             190
        4132 Leu Gly Pro Glu Phe Gly Gly Ala Val Gly Leu Cys Phe Tyr Leu Gly
        4133             195             200             205
        4135 Thr Thr Phe Ala Gly Ala Met Tyr Ile Leu Gly Thr Ile Glu Ile Phe
        4136         210             215             220
        4138 Leu Thr Tyr Ile Ser Pro Ser Ala Ala Ile Phe Gln Ala Glu Thr Ala
        4139 225             230             235             240
        4141 Asp Gly Glu Ala Ala Ala Leu Leu Asn Asn Met Arg Val Tyr Gly Ser
        4142             245             250             255
        4144 Cys Ala Leu Ala Leu Met Ala Val Val Val Phe Val Gly Val Lys Tyr
        4145             260             265             270
        4147 Val Asn Lys Leu Ala Leu Val Phe Leu Ala Cys Val Val Leu Ser Ile
        4148             275             280             285
        4150 Leu Ala Ile Tyr Ala Gly Val Ile Lys Thr Ala Phe Ala Pro Pro Asp
        4151             290             295             300
        4153 Ile Pro Val Cys Leu Leu Gly Asn Arg Thr Leu Ala Asn Arg Asn Phe
        4154 305             310             315             320
        4156 Asp Thr Cys Ala Lys Met Gln Val Val Ser Asn Gly Thr Val Thr Thr
        4157             325             330             335
        4159 Ala Leu Trp Arg Leu Phe Cys Asn Gly Ser Ser Leu Gly Ala Thr Cys
        4160             340             345             350
        4162 Asp Glu Tyr Phe Ala Gln Asn Asn Val Thr Glu Ile Gln Gly Ile Pro
        4163             355             360             365
        4165 Gly Val Ala Ser Gly Val Phe Leu Asp Asn Leu Trp Ser Thr Tyr Ser
        4166             370             375             380
        4168 Asp Lys Gly Ala Phe Val Glu Lys Lys Gly Val Ser Ser Val Pro Val
        4169 385             390             395             400
        4171 Ser Glu Glu Ser Arg Pro Gly Gly Leu Pro Tyr Val Leu Thr Asp Ile
        4172             405             410             415
        4174 Met Thr Tyr Phe Thr Met Leu Val Gly Ile Tyr Phe Pro Ser Val Thr
        4175             420             425             430
        4177 Gly Ile Met Ala Gly Ser Asn Arg Ser Gly Asp Leu Lys Asp Ala Gln
        4178             435             440             445
        4180 Lys Ser Ile Pro Thr Gly Thr Ile Leu Ala Ile Val Thr Thr Ser Phe
        4181             450             455             460
        4183 Ile Tyr Xaa Ser Cys Ile Val Leu Phe Gly Ala Cys Ile Glu Gly Val
```

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```
     4184 465                470                475                480
     4186 Val Leu Arg Asp Lys Phe Gly Glu Ala Leu Gln Gly Asn Leu Val Ile
     4187             485                490                495
     4189 Gly Met Leu Ala Trp Pro Ser Pro Trp Val Ile Val Ile Gly Ser Phe
     4190             500                505                510
     4192 Phe Ser Thr Cys Gly Ala Gly Leu Gln Ser Leu Thr Gly Ala Pro Arg
     4193             515                520                525
     4195 Leu Leu Gln Ala Ile Ala Arg Asp Gly Ile Ile Pro Phe Leu Gln Val
     4196         530                535                540
     4198 Phe Gly His Gly Lys Ala Asn Gly Glu Pro Thr Trp Ala Leu Leu Leu
     4199 545                550                555                560
     4201 Thr Ala Leu Ile Cys Glu Thr Gly Ile Leu Ile Ala Ser Leu Asp Ser
     4202             565                570                575
W--> 4204 Val Ala Pro Ile Leu Ser Met Phe Phe Xaa Met Cys Tyr Met Phe Val
     4205             580                585                590
     4207 Asn Leu Ala Cys Ala Val Gln Thr Leu Leu Arg Thr Pro Asn Trp Arg
     4208             595                600                605
     4210 Pro Arg Phe Lys Phe Tyr His Trp Thr Leu Ser Phe Leu Gly Met Ser
     4211         610                615                620
W--> 4213 Leu Cys Leu Ala Leu Met Phe Ile Cys Ser Trp Tyr Tyr Ala Xaa Phe
     4214 625                630                635                640
     4216 Ala Met Leu Ile Ala Gly Cys Ile Tyr Lys Tyr Ile Glu Tyr Arg Gly
     4217             645                650                655
     4219 Ala Glu Lys Glu Trp Gly Asp Gly Ile Arg Gly Leu Ser Leu Asn Ala
     4220             660                665                670
     4222 Ala Arg Tyr Ala Leu Leu Arg Val Glu His Gly Pro Pro His Thr Lys
     4223             675                680                685
     4225 Asn Trp Arg Pro Gln Val Leu Val Met Leu Asn Leu Asp Ser Glu Gln
     4226         690                695                700
     4228 Cys Val Lys His Pro Arg Leu Leu Ser Phe Thr Ser Gln Leu Lys Ala
     4229 705                710                715                720
     4231 Gly Lys Gly Leu Thr Ile Val Gly Ser Val Leu Glu Gly Thr Tyr Leu
     4232             725                730                735
     4234 Asp Lys His Val Glu Ala Gln Arg Ala Glu Glu Asn Ile Arg Ser Leu
     4235             740                745                750
     4237 Met Ser Ala Glu Lys Thr Lys Gly Phe Cys Gln Leu Val Val Ser Ser
     4238             755                760                765
     4240 Asn Leu Arg Asp Gly Ala Ser His Leu Ile Gln Ser Ala Gly Leu Gly
     4241         770                775                780
     4243 Gly Met Lys His Asn Thr Val Leu Met Ala Trp Pro Glu Ala Trp Lys
     4244 785                790                795                800
     4246 Glu Ala Asp Asn Pro Phe Ser Trp Lys Asn Phe Val Asp Thr Val Arg
     4247             805                810                815
     4249 Asp Thr Thr Ala Ala His Gln Ala Leu Leu Val Ala Lys Asn Ile Asp
     4250             820                825                830
     4252 Leu Phe Pro Gln Asn Gln Glu Arg Phe Ser Asp Gly Asn Ile Asp Val
     4253             835                840                845
     4255 Trp Trp Ile Val His Asp Gly Gly Met Leu Met Leu Leu Pro Phe Leu
     4256         850                855                860
```

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

```
      4258 Leu Arg Gln His Lys Val Trp Arg Lys Cys Arg Met Arg Ile Phe Thr
      4259 865               870               875               880
      4261 Val Ala Gln Val Asp Asp Asn Ser Ile Gln Met Lys Lys Asp Leu Gln
      4262               885               890               895
      4264 Met Phe Leu Tyr His Leu Arg Ile Ser Ala Glu Val Glu Val Val Glu
      4265               900               905               910
      4267 Met Val Glu Asn Asp Ile Ser Ala Phe Thr Tyr Glu Lys Thr Leu Met
      4268               915               920               925
      4270 Met Glu Gln Arg Ser Gln Met Leu Lys Gln Met Gln Leu Ser Lys Asn
      4271         930               935               940
      4273 Glu Arg Glu Arg Glu Ala Gln Leu Ile His Asp Arg Asn Thr Ala Ser
      4274 945               950               955               960
      4276 His Thr Thr Ala Thr Ala Arg Thr Gln Ala Pro Pro Thr Pro Asp Lys
      4277               965               970               975
      4279 Val Gln Met Thr Trp Thr Lys Glu Lys Leu Ile Ala Glu Lys His Arg
      4280               980               985               990
      4282 Asn Lys Asp Thr Gly Pro Ser Gly Phe Lys Asp Leu Phe Ser Leu Lys
      4283               995               1000              1005
W--> 4285 Pro Asp Gln Ser Asn Val Arg Arg Met His Thr Ala Val Lys (Xaa) Asn
      4286       1010              1015              1020
      4288 Gly Val Val Leu Asn Lys (Ser) Gln Asp Ala Gln (Leu) Val Leu Leu Asn
E--> 4289 1025              ∠1030              ∠1035             ∠1040        see below
      4291 Met Pro Gly Pro Pro Lys Ser Arg Gln Gly Asp Glu Asn Tyr Met Glu
      4292               1045              1050              1055
      4294 Phe Leu Glu Val Leu Thr Glu Gly Leu Asn Arg Val Leu Leu Val Arg
E--> 4295       1060              1065              1070
      4297 Gly Gly Gly Arg Glu Val Ile Thr Ile Tyr Ser
      4298         1075              1080
```

1) begin the amino acid number directly below the first letter of the amino acid. Do not have any part of the number under the letters of the next amino acid,

e.g. Lys | s p a c e | Ser
     1030

2). When numbering the last amino acid on a line, please end the number directly below the last letter of the amino acid
   Leu | s p a c e | Asn
       1040

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```
L:14   M:270 C: Current Application Number differs, Replaced Application Number
L:15   M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:59   M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:60   M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:67   M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:68   M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:303  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:304  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:417  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:423  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:600  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:650  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:651  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:762  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:763  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:910  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:911  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:981  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:1062 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:1176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:1239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1240 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1630 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:1729 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:1819 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:1858 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:1859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:1926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:1927 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2062 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2063 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2630 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2631 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2739 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:2895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:2982 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:3063 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:3143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:3144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:3203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:3204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
```

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Input Set : **A:\EP.txt**
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```
L:3343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:3344 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:3524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:3569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:4289 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14
M:332 Repeated in SeqNo=14
L:6706 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:111
L:6834 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:112
L:7185 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:131
```